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FKMPKMTMPKLGRAESPSRGKPGEAGAEVSGKLVTLPGEDEARVGVPSLTLPS
VELDLPGALGLOGOVPAAKMGKGERVEGKLVTLPGFFRVPSVEIVTPOLPAVEIE
EGRLEMIETKVKPSSKFSLPKFGLSGPKVAKAEAEGAGRATKLKVSKFAISLPKARVG
RAEAEAKGAGEAGLLPALDLSIPOLSLDAHLPSGKVEVAGADLKFKGPRFALFKFGVRG
RDTEAAELVPGVAELEGKGWGNDGRVKMPKLKMPSFGLARGKEAEVQGDRASPGEKAE
STAVQLKIPEVELVTLGAQEEGRAEGAVAVSGKQLSGLKVSTARQVVTEGEDAGLRMF
STAVQLKIPEVELVTLGAGAEGCRAEGAVAVSGKDLSGLKVSTARQVVTEGEDAGLRMF
PLGISLPGVELTGFGEAGTFGQQAQSTVPSAEGTAGYRVQLVGVTLGLFGAFARVGGEGAE
LVGEGVFKMPTVTVPQLELDVGLSFEAGAGFAGVAGGEGLRKLKVRLPREGLVRAKEGA
EGGEKAKSPKLKLPRVGFSGSMVTGBGSPSBEEEEBEEEGGAGSGRGRGRVRVRL
EGGEKKASPKLKLPRVGFSGSMVTGBGSPSBEEEEBEEEGGAGAGRGRGRVRVRL
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alternatively spliced"
/codon_start=1
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270. .4655
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/protein_id="AAK19279.1"
/db_xref="GI:13345275"
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="19"
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1. .4853
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601 CCGTGTCTGGCTACGAGATCAAGGGCCCGCGGGCCAAGGTGGCCAAGCTGAACATCCAGA
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tive 0; Mismatches
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| Qу | Qy Db | D Qy | Qy Db | Qу | Qy | dg Vo | p Q | Db Qy | do Qy | Qy Db | Qy Db | Qy Db | Qy | Оу | Оу | Qy | Оy | Db |
|--|--|---|--|--|--|---|--|--|---|--|--|--|--|---|--|---|--|--|
| 8=8 | 1801 CTGTGCCGGAGGTGCGGCTTCCAGAGGTACAGCTGCTGAAAGTGTCGGAGATGAAACTCC 1860 | 1741 TTCCAGAGGTAGAGCTGCCCAAAGTGTCAGAGATGAAACTCCCAAAAGGTGCCAGAGATGG 1800 | 1681 CCAAGGTGTCAGAGATGAAACTCCCCAAAGGTGCCAGAGATGGCTGTGCCGGAGGTGCGGC 1740 | 1621 AGCTTCCAAAAGTGCCCGAGGCAGCCCTTCCAGAGGTTCGACTCCCAGAGGTGGAGCTCC 1680 1611 1 | 1561 GGCCCGAGGTCAAGGTGCCCAAGGGACCTGAAGTGAAGCTCCCCAAGGCTCCTGAGGTCA 1620 | 1501 TAGAGAGCAAGCTGAAGCTGCCCACCATCAAGATGCCCTCCCT | 1441 TTCGAATGCCCACCITTGGGCITTCCCTCTTGGAGCCCCGGCCCGCTGCTCCTGAAGTTG 1500 | 1381 AAGTTGCTGAGGCCAAGGTAGCCAAGGTCAGCCCTGAGGCCAGGGTGAAAGGTCCCCAGAC 1440 | 1321 CTGAGGTGGCCCTGAAGATGCCCCGCCTTAGTTTTCCCCCGATTTGGGGCTCGAGCAAAGG 1380 | 1261 CGACTGTGGGGGTGGACCTGGCCTTGCCGGGTGCAGACGTGGAGGCCCCGGGGAGAGGCAC 1320 | 1201 GCCTAGAGACCCGGGAAGGGGCTGTGTCGGTAGTGGTGCCCACCCTGGATGTGGCAGCAC 1260 | 1141 AGGTCCCCCAGGTGGAGCTGCCTGCCCTGCCCTCACTGCCCACACTTCCCT 1200 | 1081 CAACCCTTGGGCTCGGAGCCCCGGCTCCGCCTGTGGAGGCCCAGCCGTGGGAATCC 1140 | 1021 TCTCAGCCCCCAAGGCTGCCCCCTCAGCAGAAGGCAGCTGGTTGGCCTTTGCCCTCCACCTGC 1080 | 961 CCCCTCAGGTGGAGCTGGTTGGGCCGCGGCGCCAGGGGCGGAGGTGGGTG | 901 CTCCTCCCCCAGGAAAGCCAAGGTGGAGGTGGAGGTGGCTGCAGGAGCTCGTTTCACAG 960 | 841 CTCGGCTGCTGTACGAGAAGTGGCCGAAGAGGCTCAGGCAGCCCGGCTGGCCGCCGCCCG 900 | 81 / |
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| 4081 CCTTCTGCCCTCTCACTGCCCACGGGAGGAGGAGGAGGAGGAGGACCACAGGCAACCATGCCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGA | ₽ . | 0 V | ф | Qy | Ф | Qy | Дb | Qy | Db | Qy | D | Qy | Дb | Qy | dd | Оу | Дb | Qγ | DЪ | Qy | Db | γQ | DЪ | VQ | Db |
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Homo sapiens Eukaryota; Metazoa; Mammalia; Eutheria; Chen,S.Y., Macina,R.A., Sun,Y. and Recipon,H. Compositions and methods relating to lung speratent: WO 0218776-A 5 07-MAR-2002; Chordata;
Primates;

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

specific

genes

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linear

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/organism="Homo sapiens" /db_xref="taxon:9606" 1660 c 1980 g 96 Location/Qualifiers .5746

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